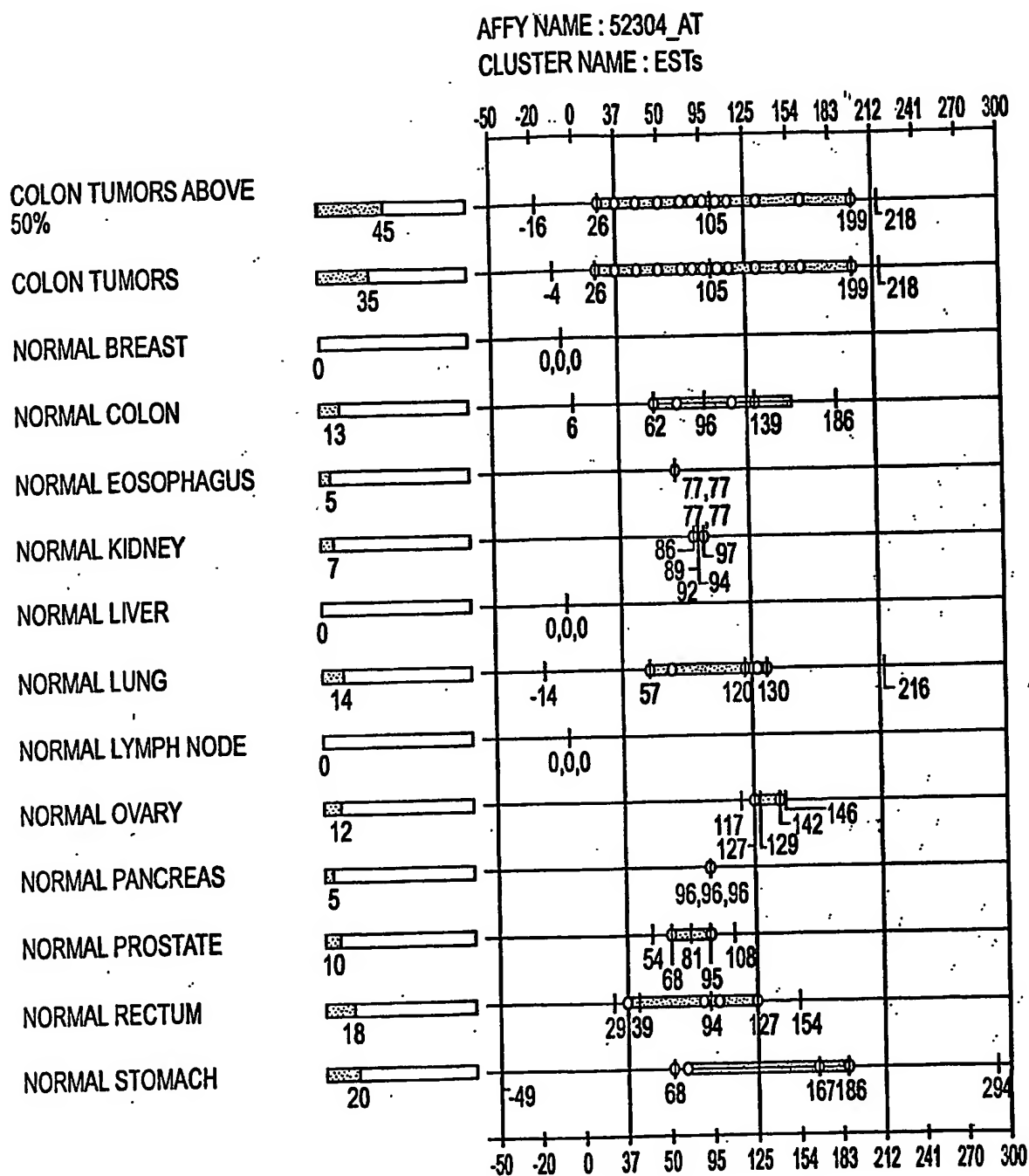
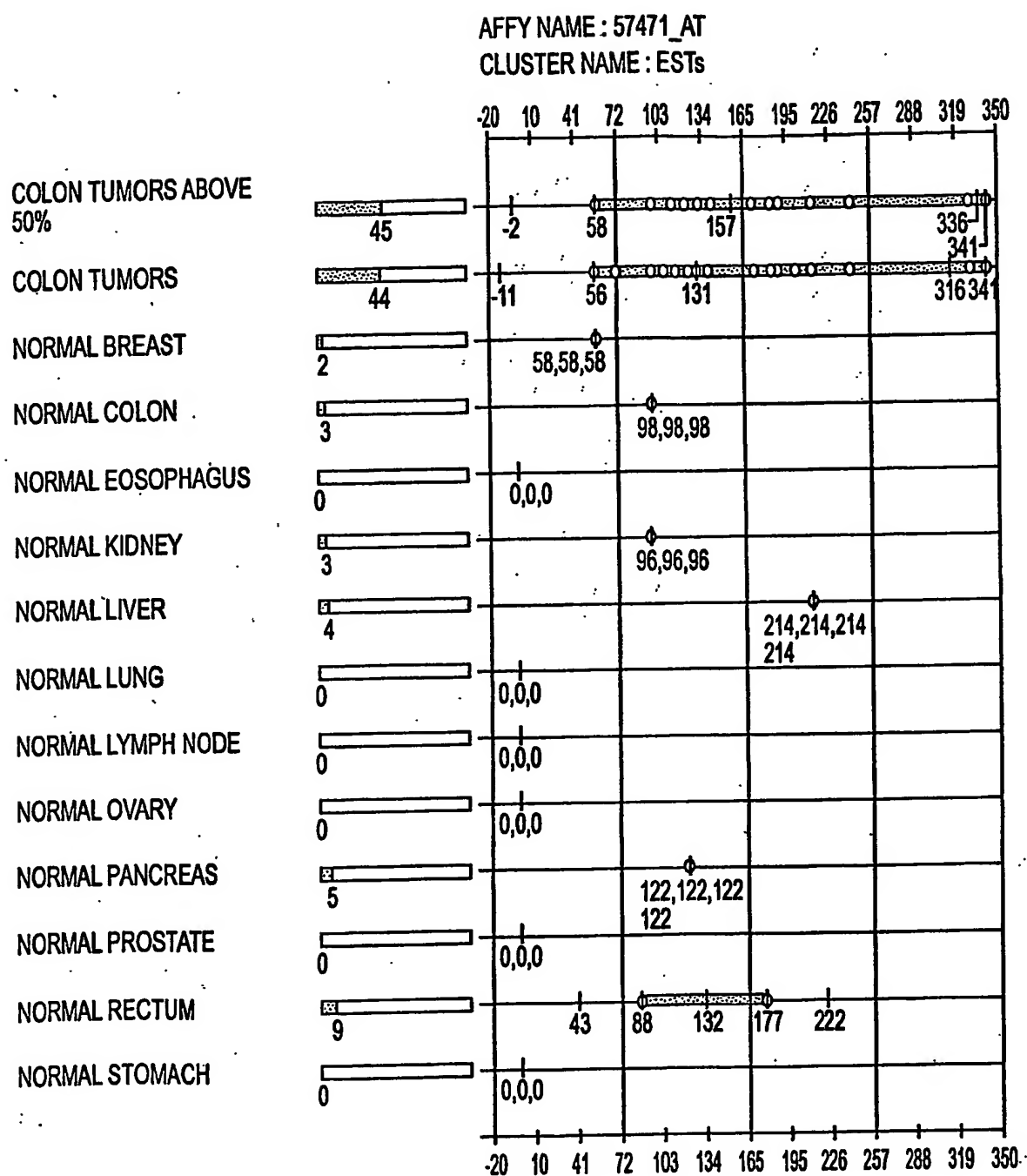
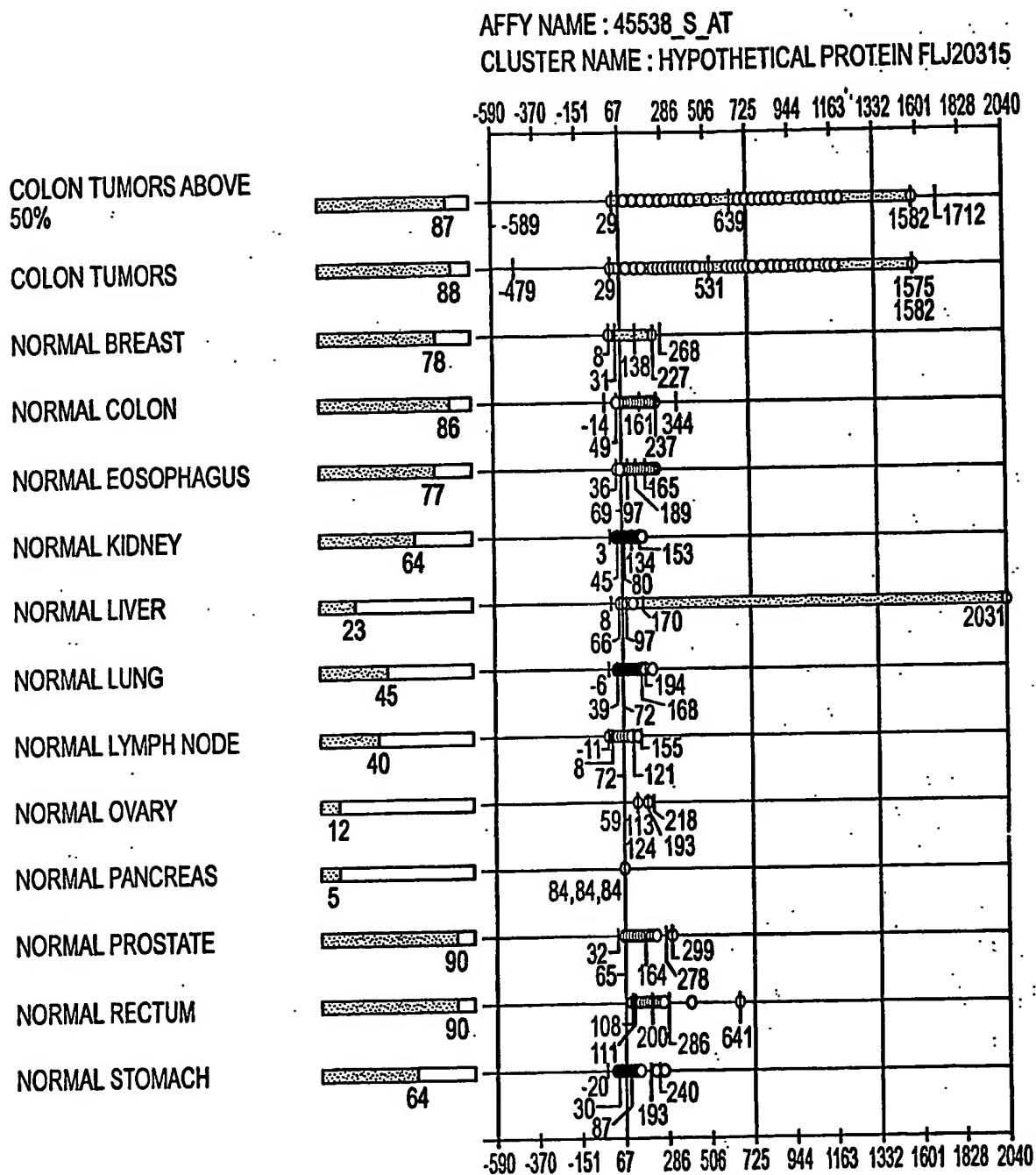


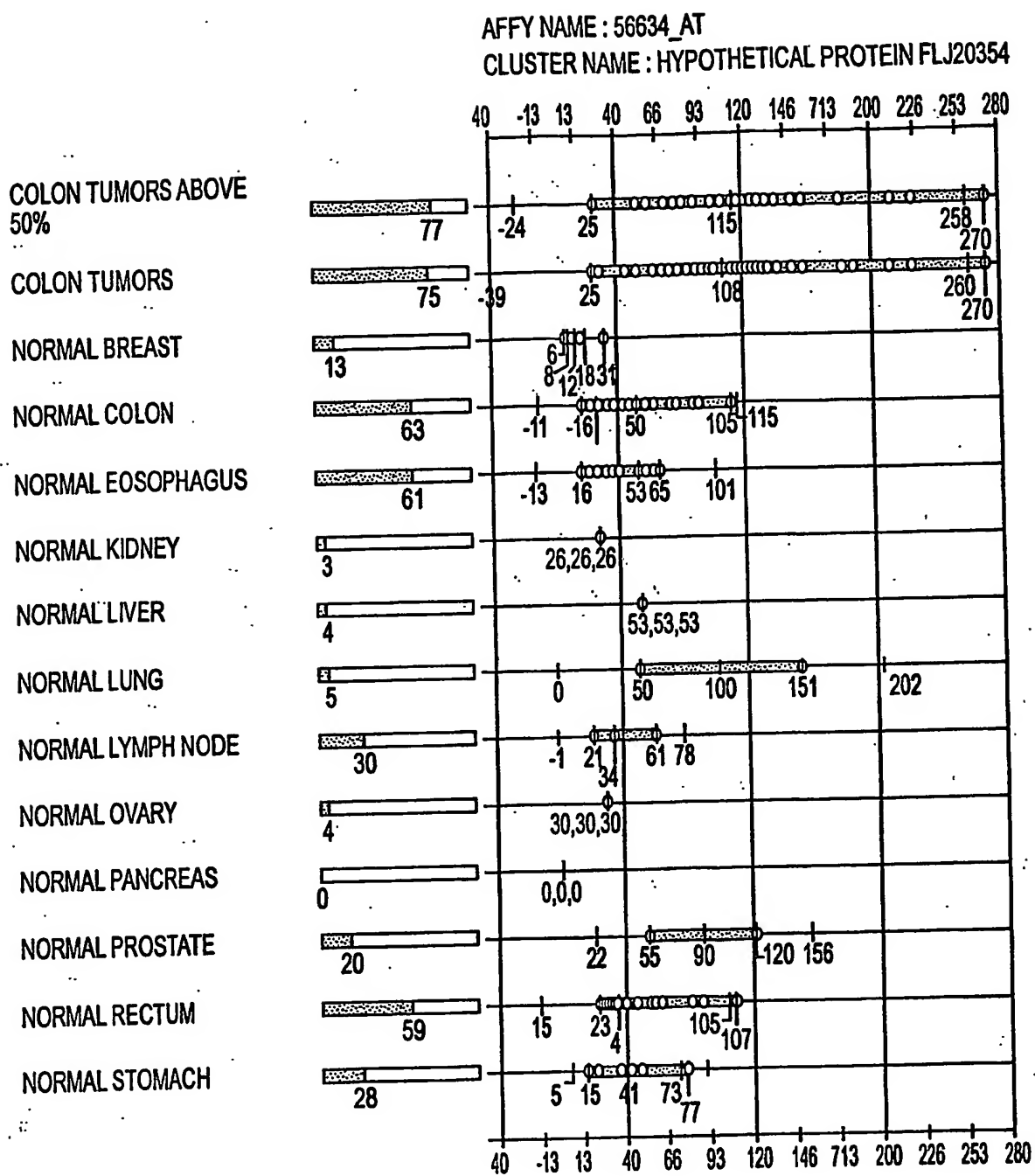
NAME	DESIGNATION	SIZE	A TUMOR	A TUMOR	AN:N	ANORMAL	BT:N	B TUMOR	POOL	POOLED NORMAL	DT:N	D TUMOR	
CICO1	BS210MS130	184.75	3	1213.04	3	256.05	3	2685.32	2	298.09	3	1812.48	
CICO2	BS220MS230	191.20	3	994.12	3	108.37	3	228.49	3	60.10	3	1271.98	
CICO3	BS430MS430	222.32	3	1587.80	3	113.79	3	1226.02	3	117.06	3	163.29	
	BS421MS433	258.00	3	213.22	3	62.02	3	175.93	2	36.16	3	158.55	
FT2:N	F TUMOR	FN2:N	F NORMAL	GT:N	G TUMOR	GN:N	G NORMAL	HT:N	H TUMOR	HN:N	H NORMAL	IT:N	I TUMOR
3	658.97	3	229.01	3	2750.16	3	307.80	3	858.10	3	271.62	3	2501.17
3	370.06	3	41.80	3	590.73	3	56.69	3	295.42	3	90.93	3	1164.92
3	2747.40	3	118.80	3	1559.78	3	112.18	3	302.67	3	66.51	3	8010.81
3	148.52	3	45.92	3	306.88	3	115.65	3	168.77	3	75.19	3	190.42
IN:N	I NORMAL	MT:N	M TUMOR	MIN:N	M NORMAL	QT:N	Q TUMOR	QN:N	Q NORMAL	RT:N	R TUMOR	RN:N	R NORMAL
3	225.34	3	3354.80	3	267.40	3	1494.28	3	261.62	3	2518.73	3	550.77
3	57.46	3	90.37	3	47.53	3	132.73	2	66.31	3	376.15	3	55.26
3	90.56	2	280.45	3	245.67	3	2593.52	3	158.56	3	1061.67	3	70.30
3	29.94	3	95.06	2	34.97	3	237.60	3	24.70	3	293.87	3	53.45
8TN	8 TUMOR	TUMOR AVE	NORMAL AVE	AVE FOLD	# OF HITS								
3	256.60	1827.61	296.41	6.17	20								
1	130.50	513.22	64.94	7.90	35								
3	160.96	1790.40	121.49	14.74	19								
3	248.80	203.42	53.11	3.83	22								

FIG. 1

**FIG. 2**

**FIG. 3**

**FIG. 4**

**FIG. 5**

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18

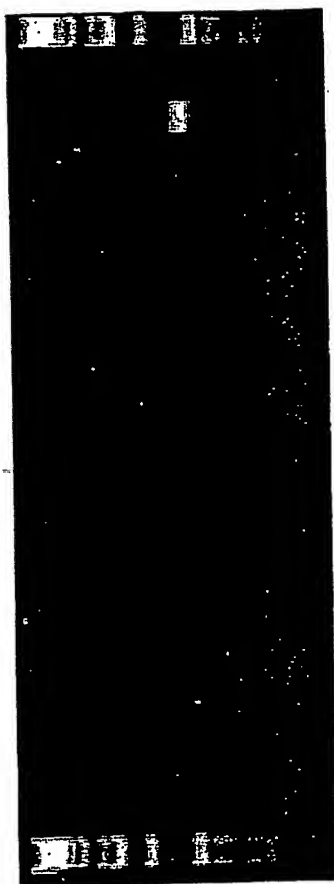


FIG. 6A

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18

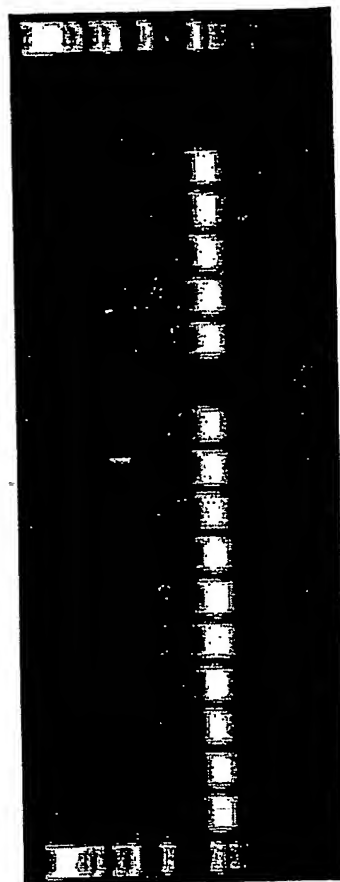


FIG. 6B

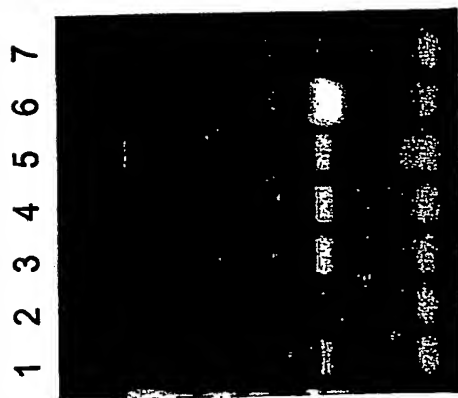


FIG. 7A

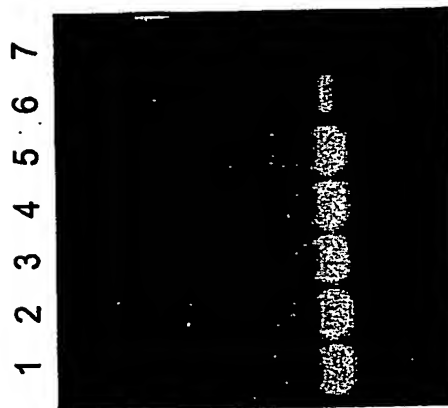
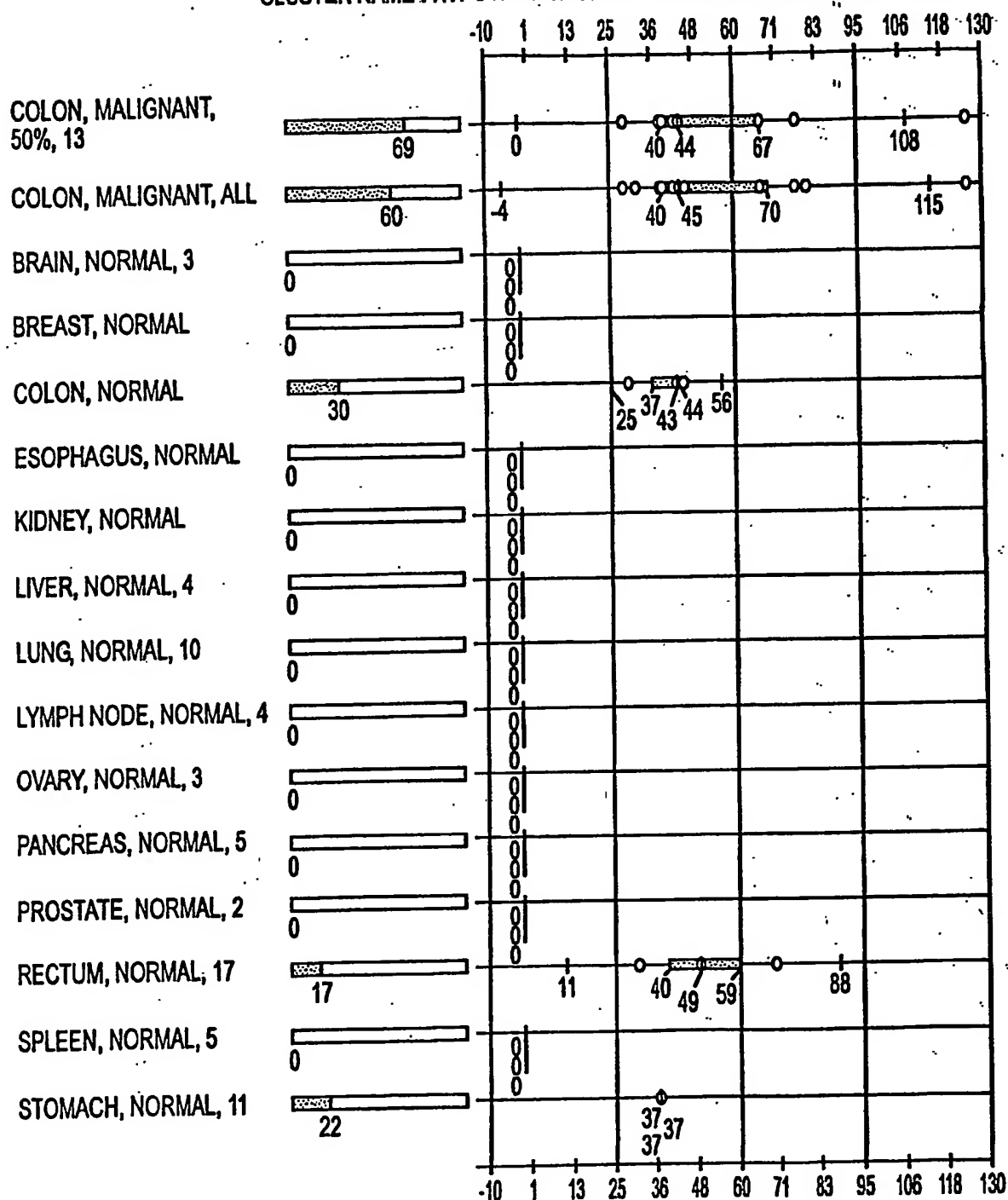


FIG. 7B

AFFY NAME : 222206_S_AT

CLUSTER NAME : HYPOTHETICAL PROTEIN FROM EUROIMAGE 2021883

**FIG. 8**

M 1 2 3 4 5 6 7 8 9 10 11 12

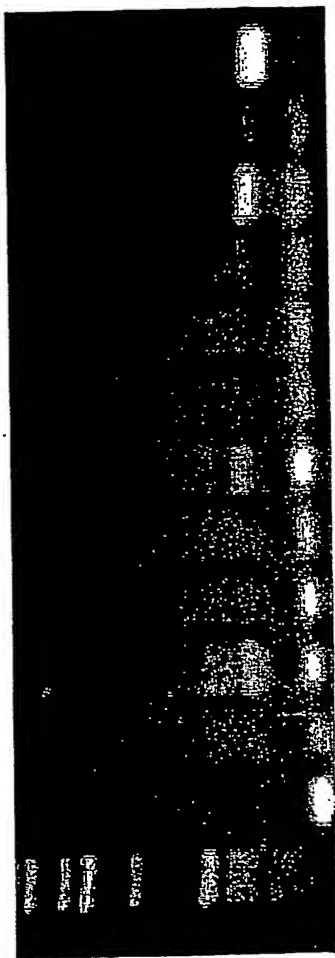


FIG. 9A

M 1 2 3 4 5 6 7 8 9 10 11 12

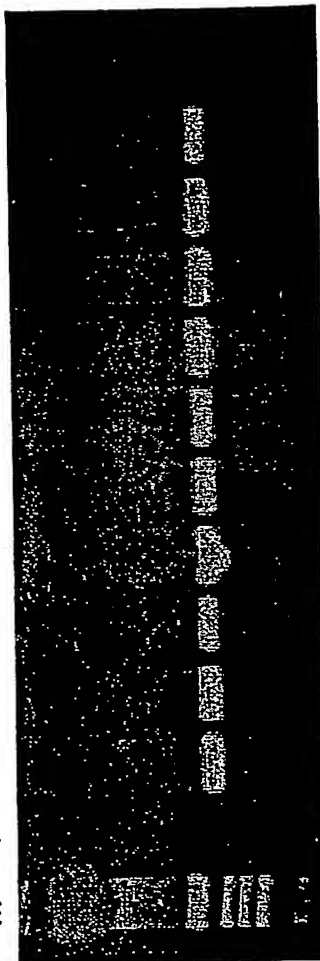


FIG. 9B

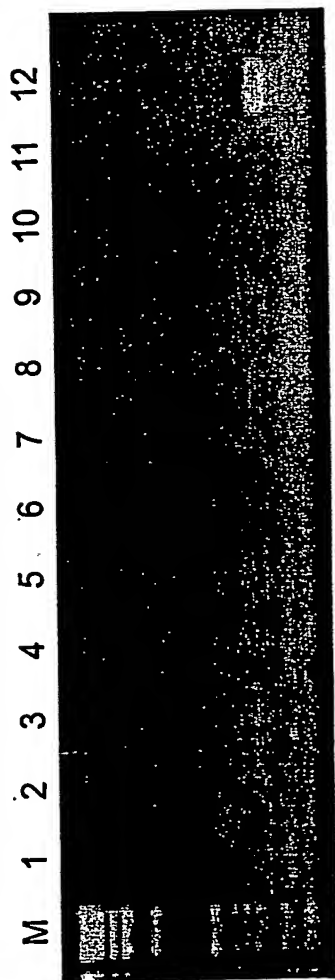


FIG. 10A

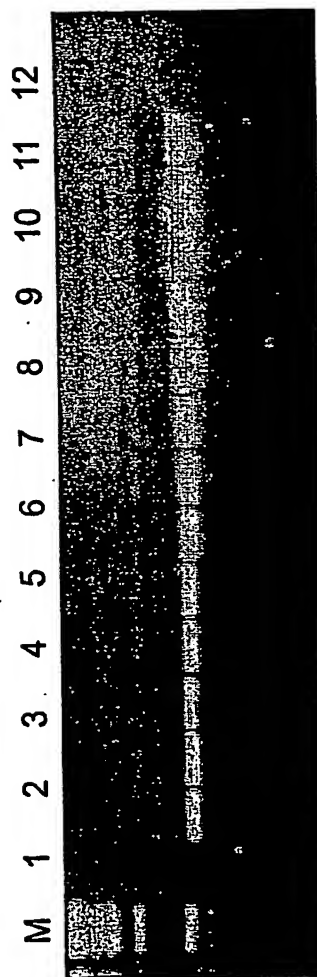


FIG. 10B

M 1 2 3 4 5 6 7 8 9 10 11 12 13 14

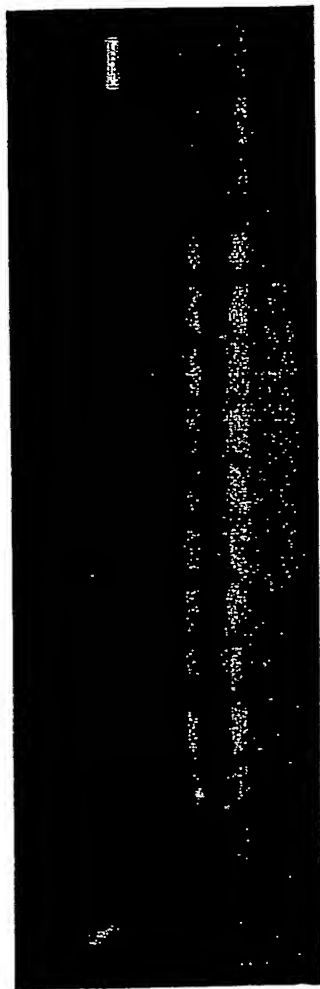


FIG. 11A

M 1 2 3 4 5 6 7 8 9 10 11 12 13 14

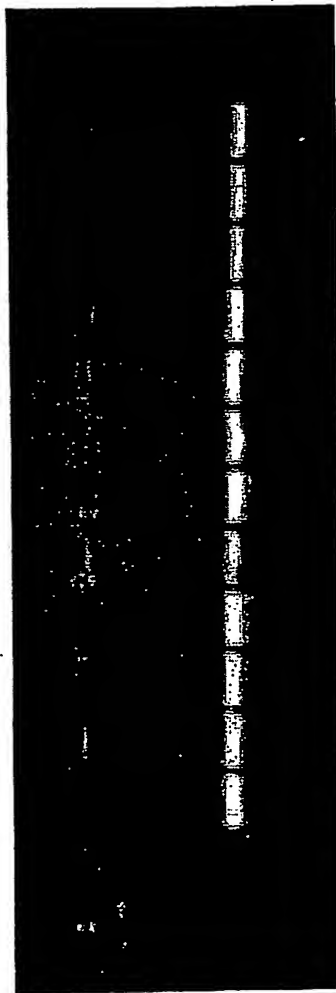


FIG. 11B

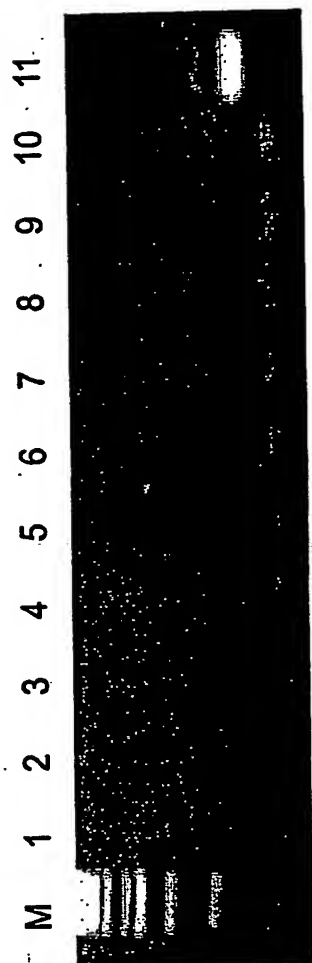


FIG. 12A

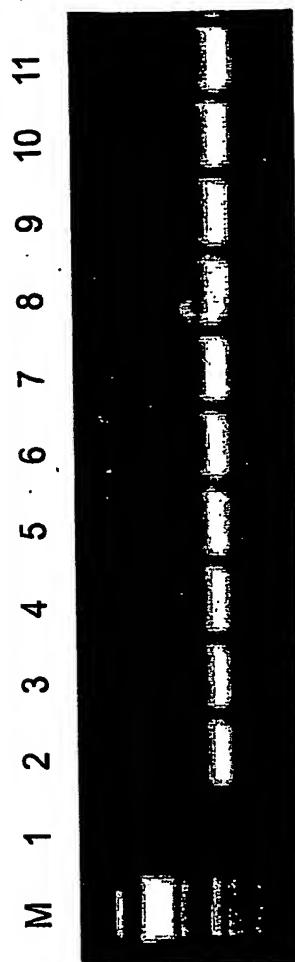
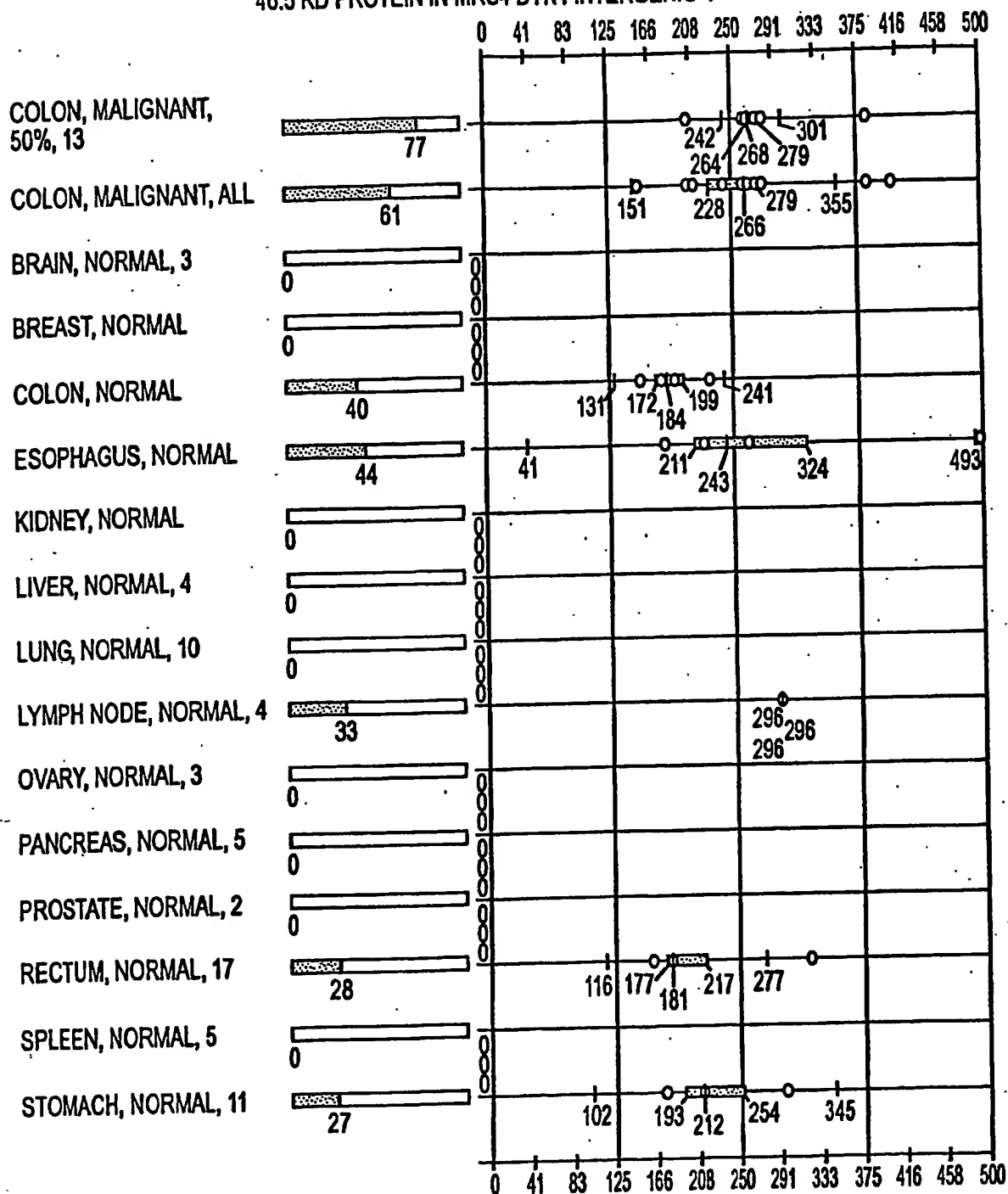


FIG. 12B

AFFY NAME : 238567_AT

CLUSTER NAME : ESTs, WEAKLY SIMILAR TO YK33_YEAST HYPOTHETICAL
46.5 KD PROTEIN IN MRS4-DYN1 INTERGENIC F**FIG. 13**

AFFY NAME : 225762_AT
 CLUSTER NAME : ESTs, WEAKLY SIMILAR TO A2AA RAT ALPHA-2A
 ADRENERGIC RECEPTOR [R.NORVEGICUS]

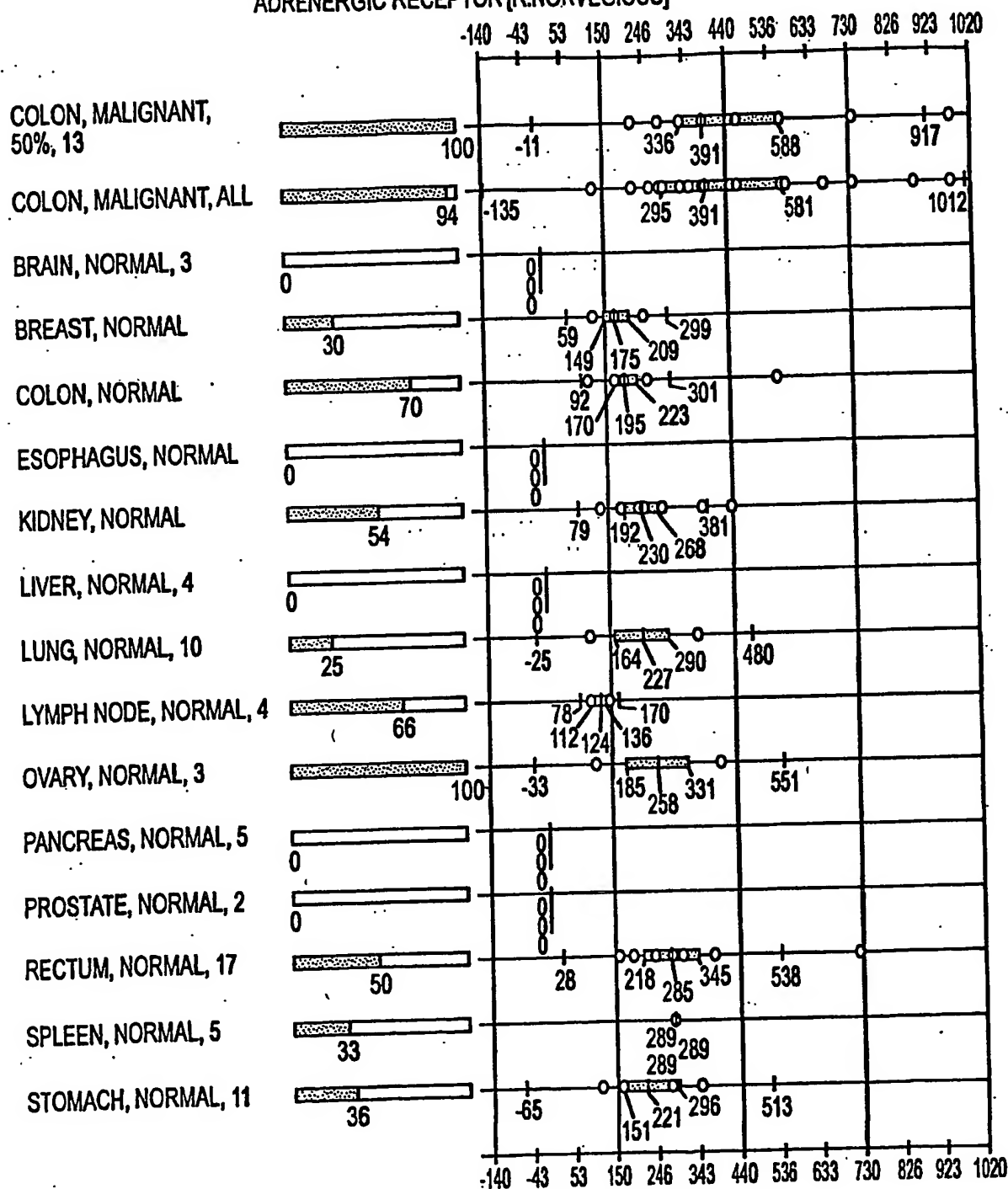
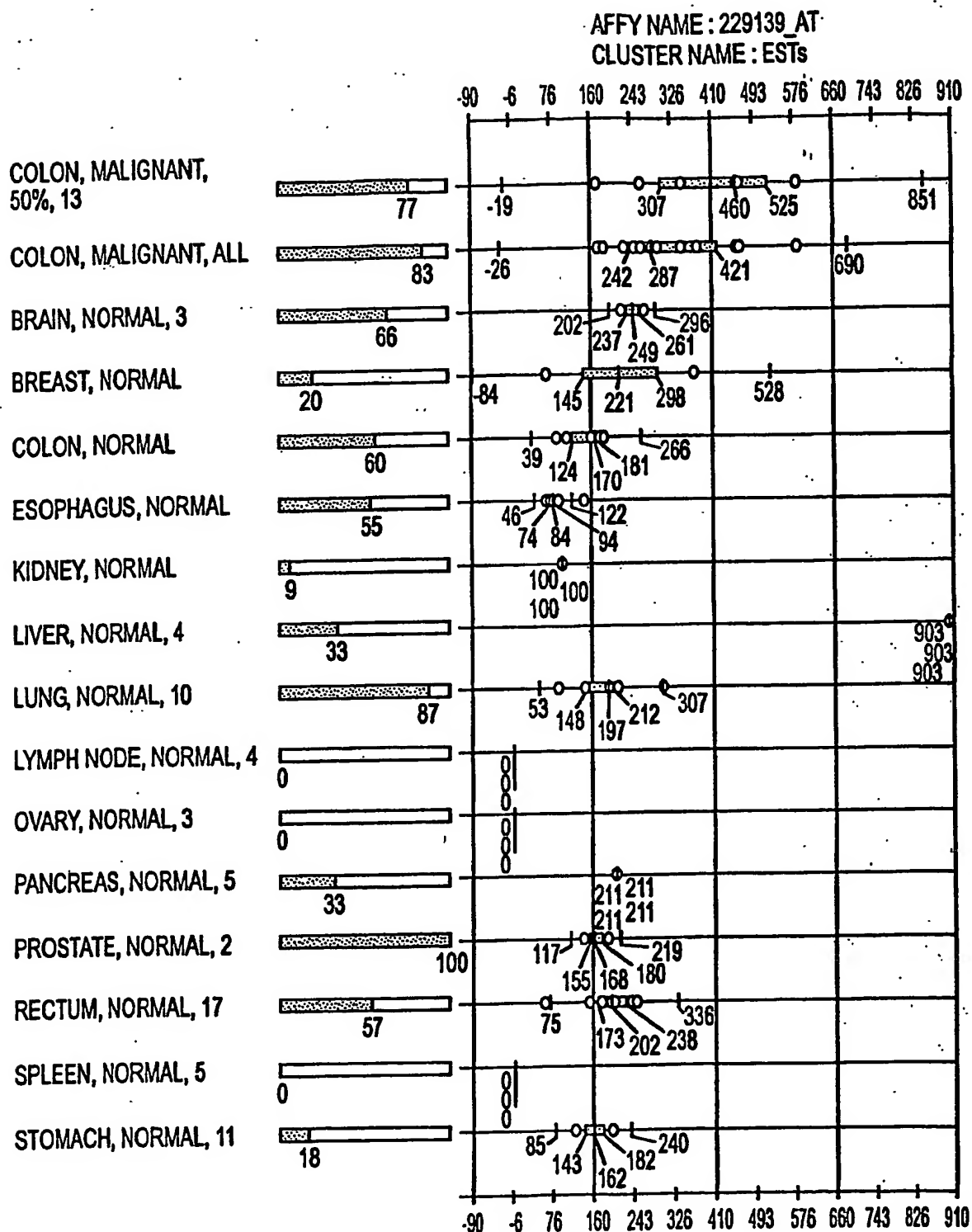


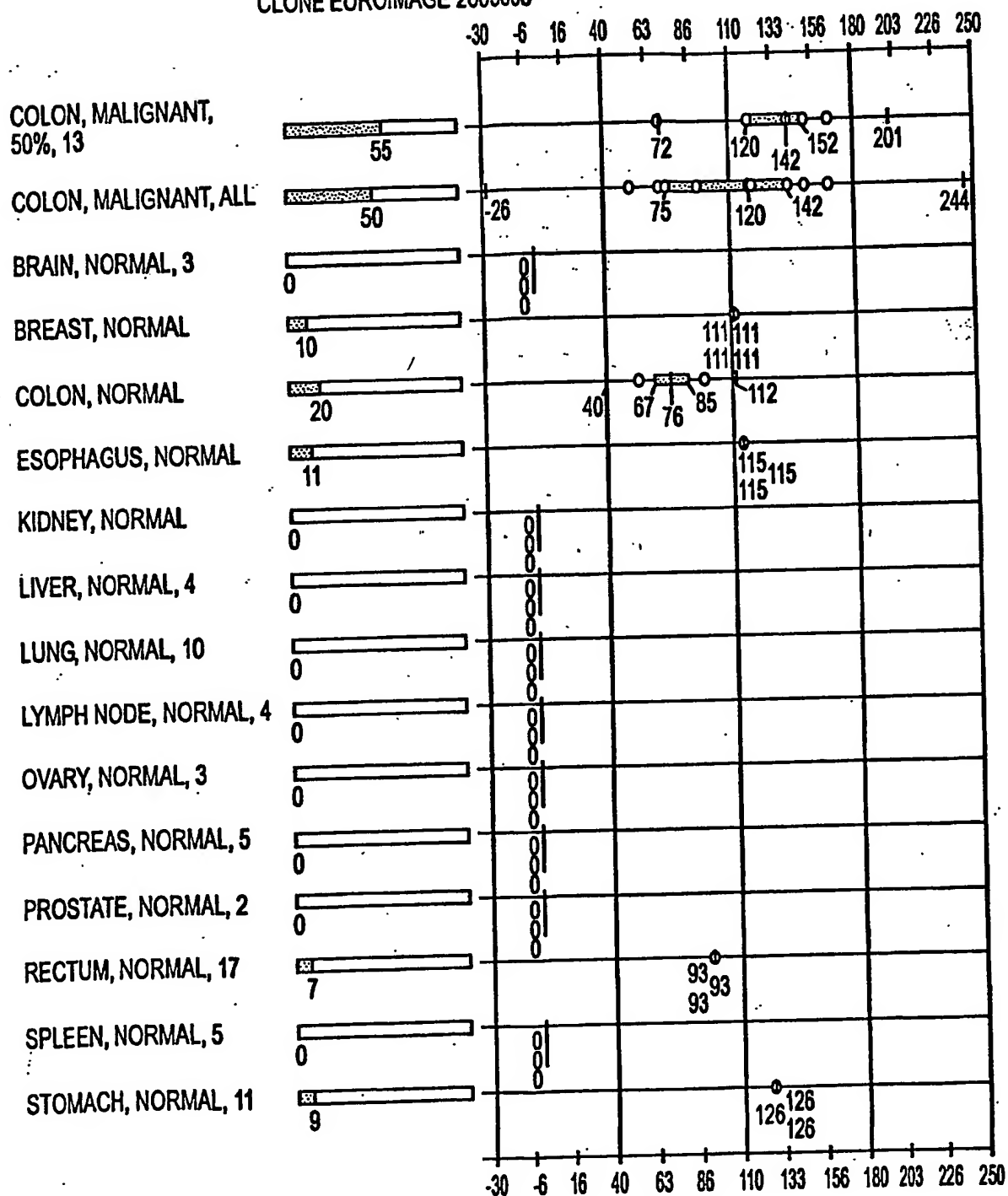
FIG. 14

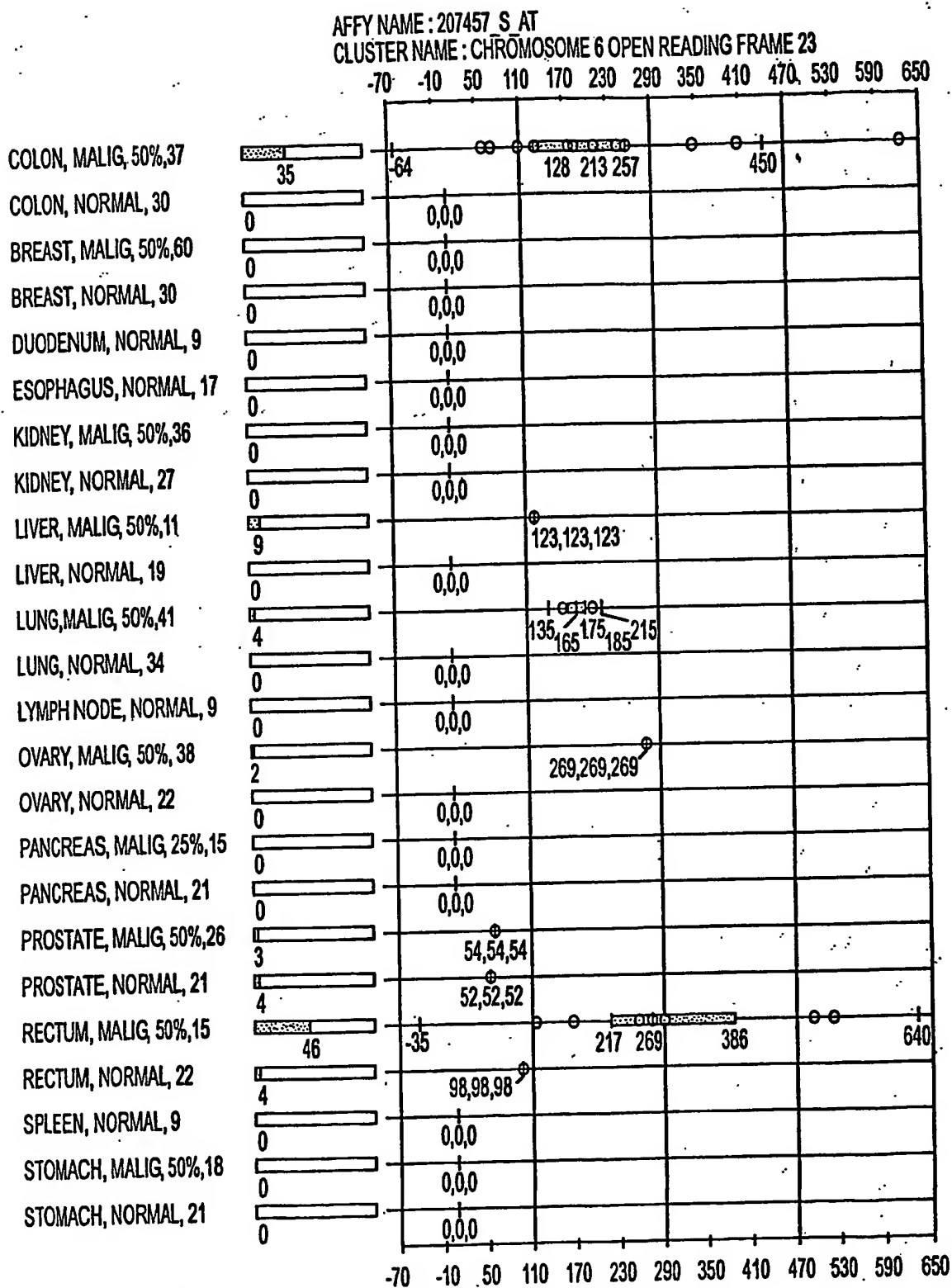
**FIG. 15**

AFFY NAME : 232504_AT

CLUSTER NAME : HOMO SAPIENS mRNA FULL LENGTH INSERT cDNA

CLONE EUROIMAGE 2005635

**FIG. 16**

**FIG. 17**

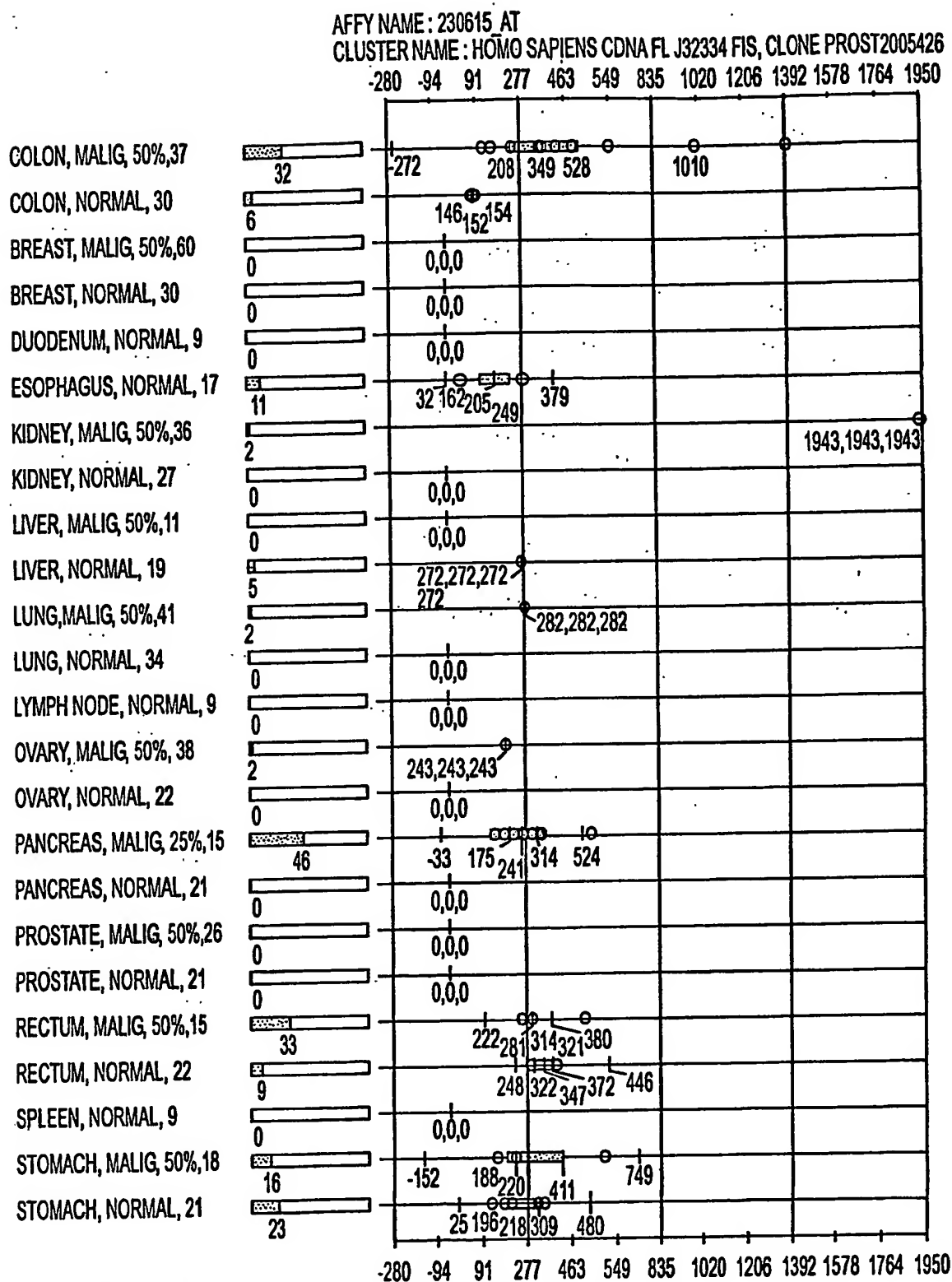
**FIG. 18**

FIG. 20A

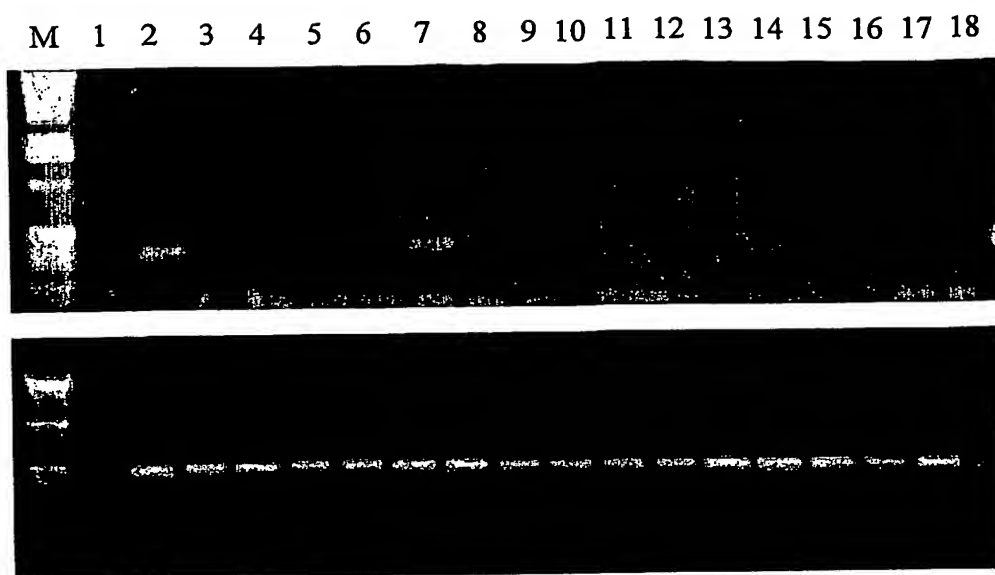


FIG. 20B

FIG. 21A

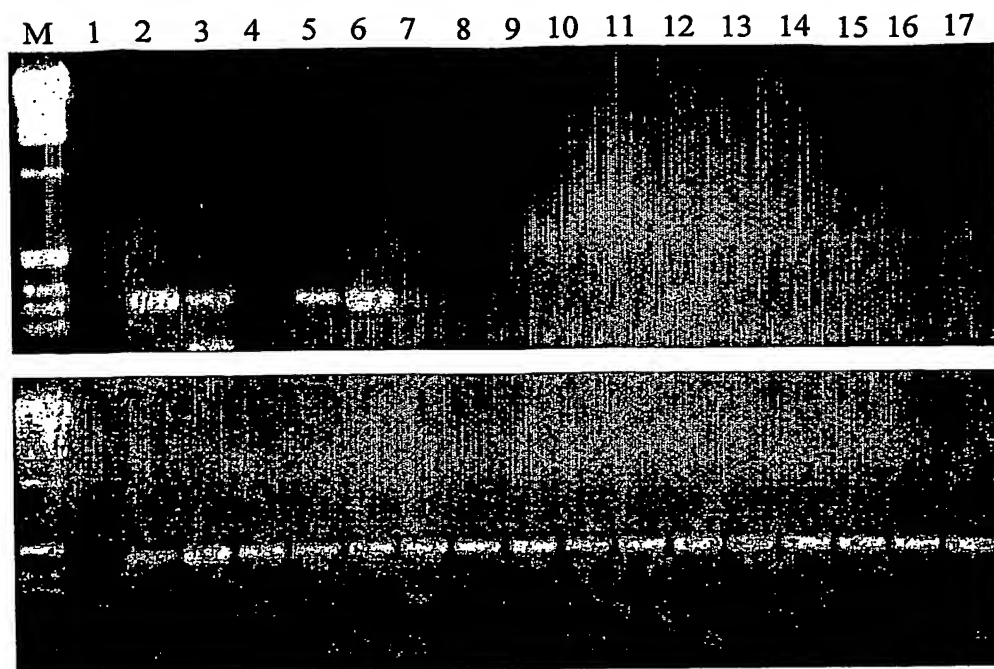


FIG. 21B

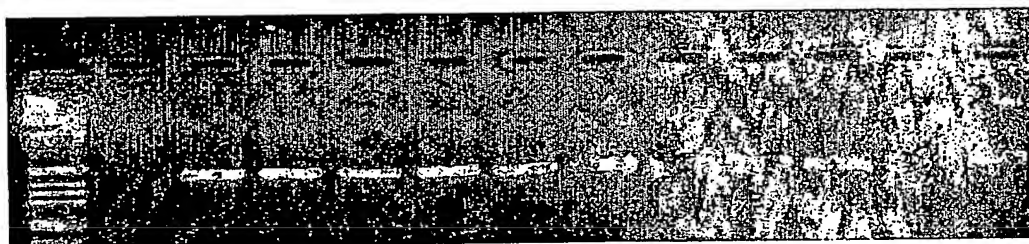
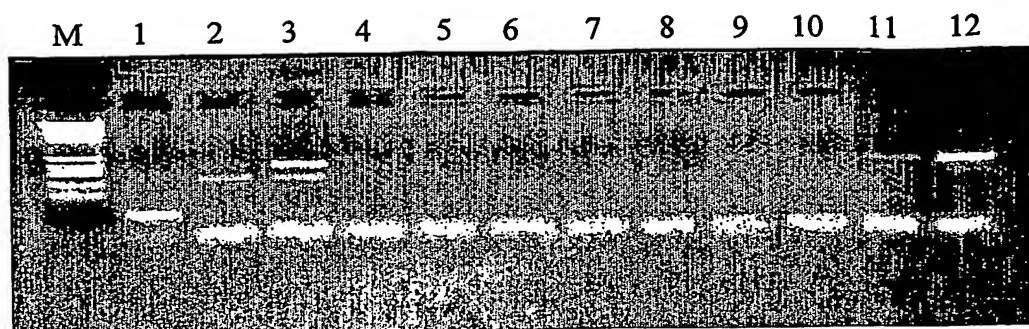
FIG. 22A*FIG. 22B*

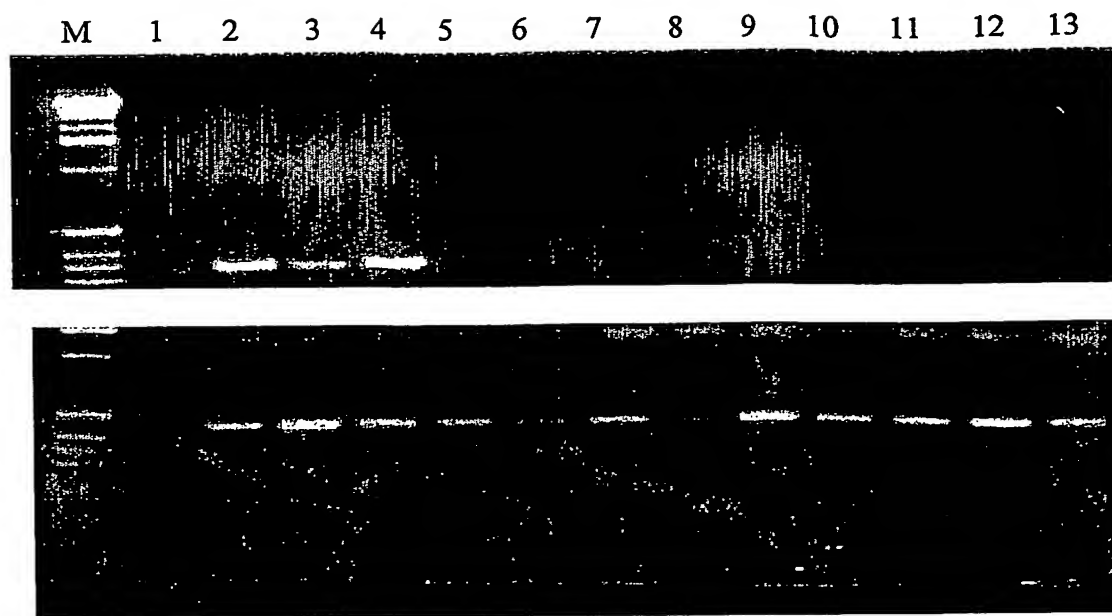
FIG. 23A*FIG. 23B*

FIG. 24A

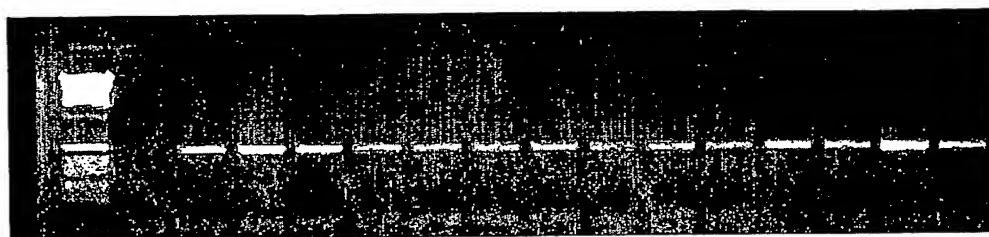
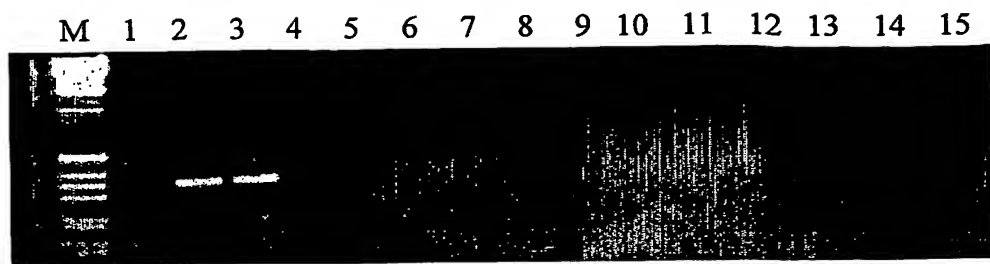


FIG. 24B

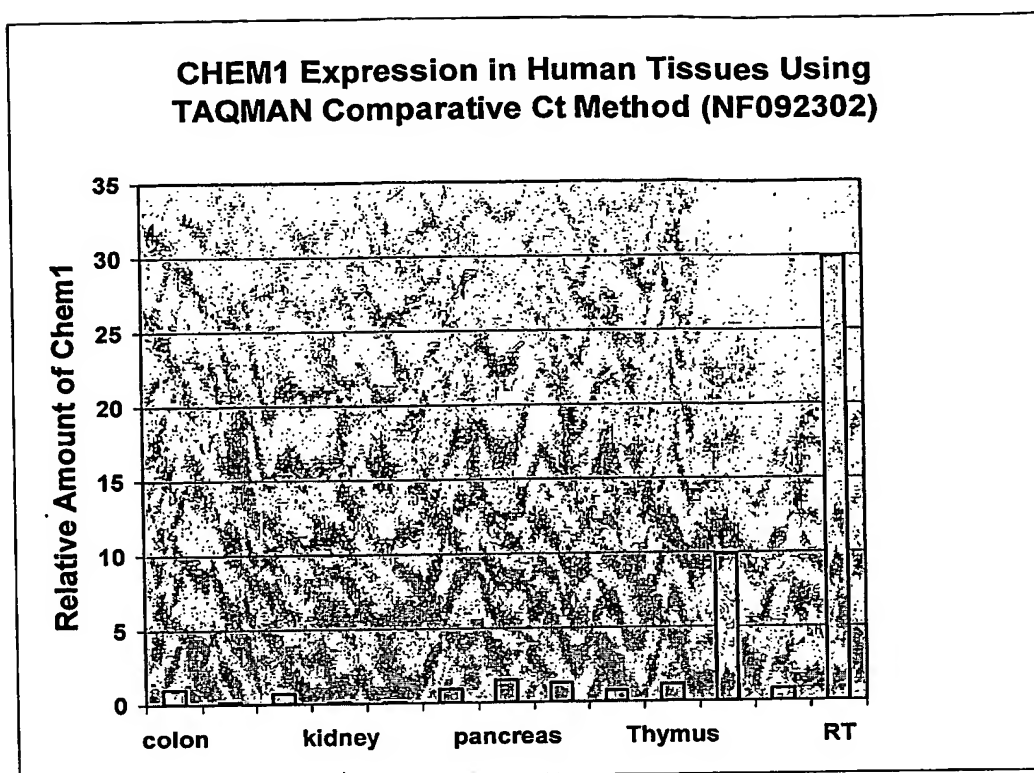
FIG. 25

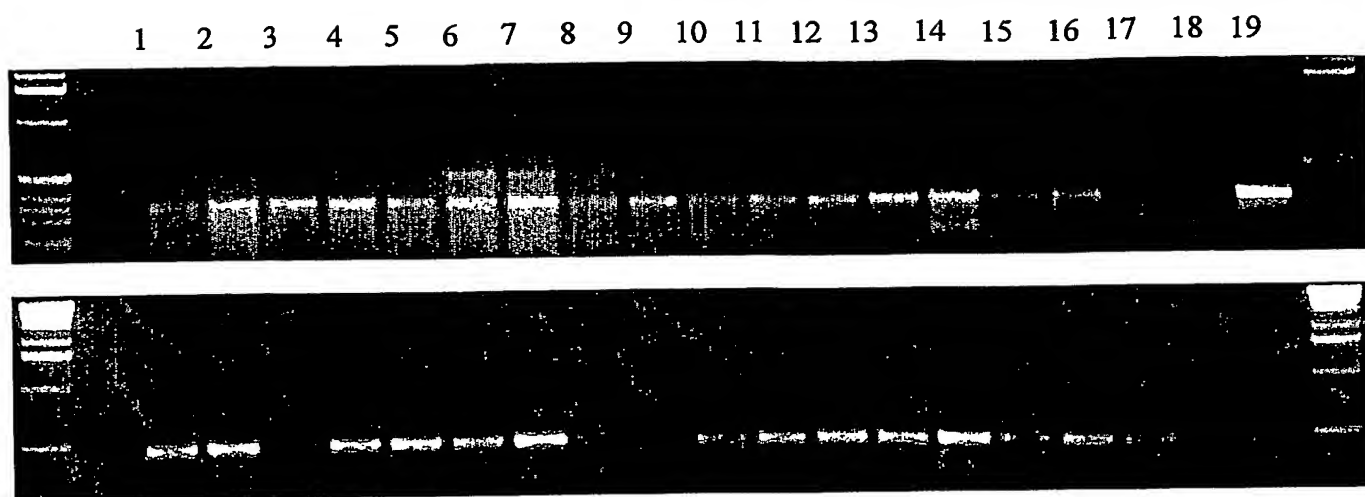
FIG. 26A*FIG. 26B*

FIG. 27

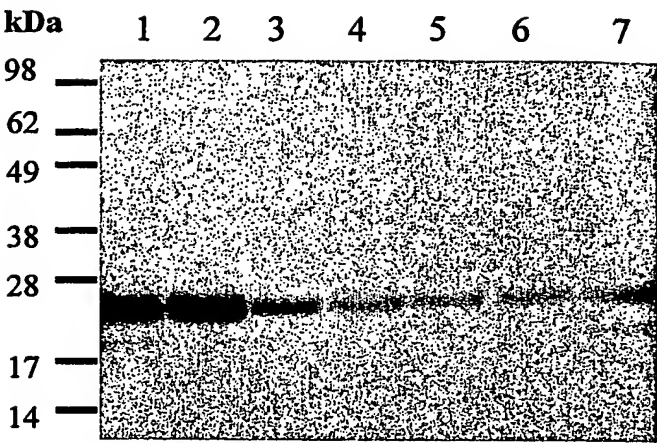


FIG. 28

